

00050" 990E3560

000E50" 990E8560

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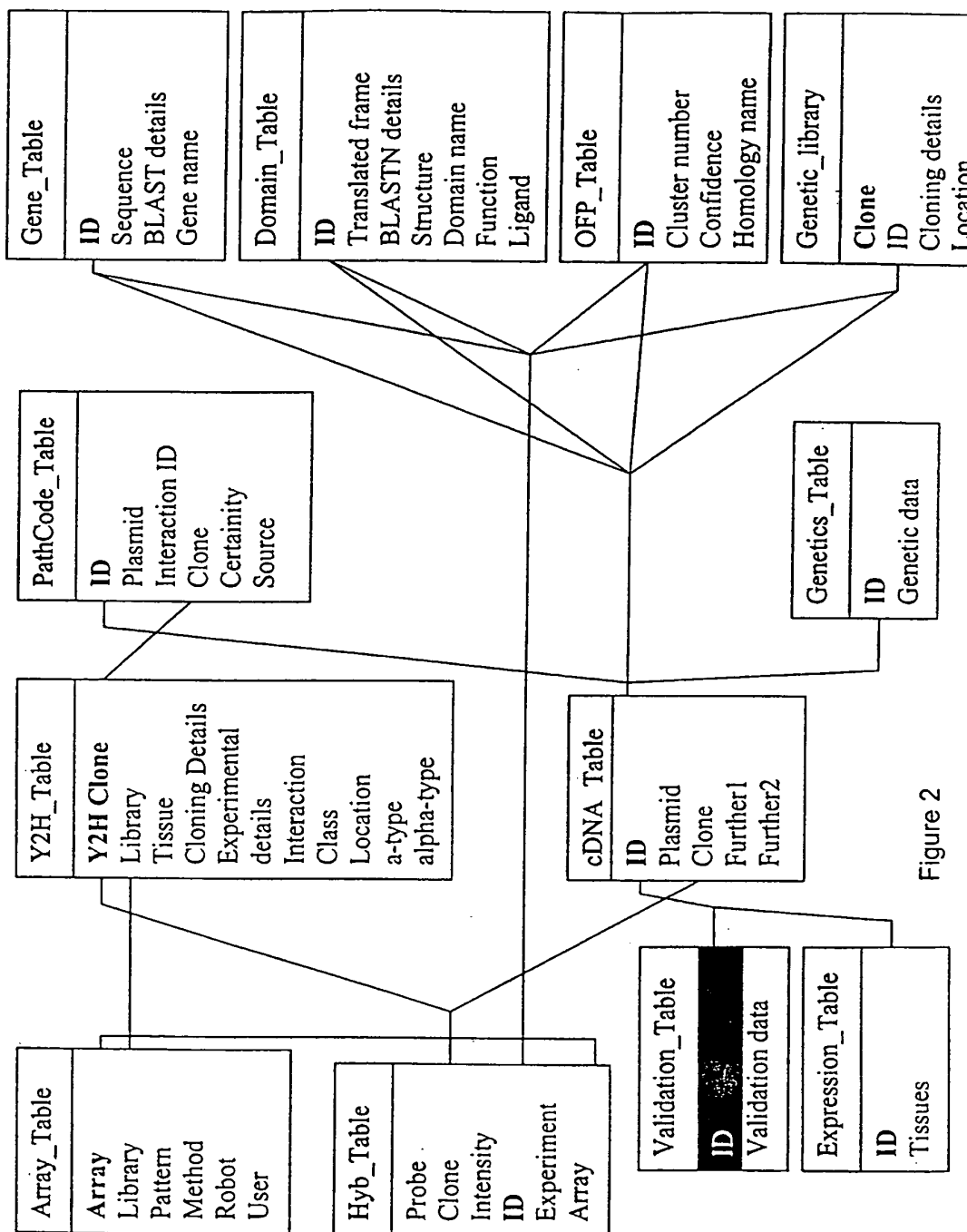
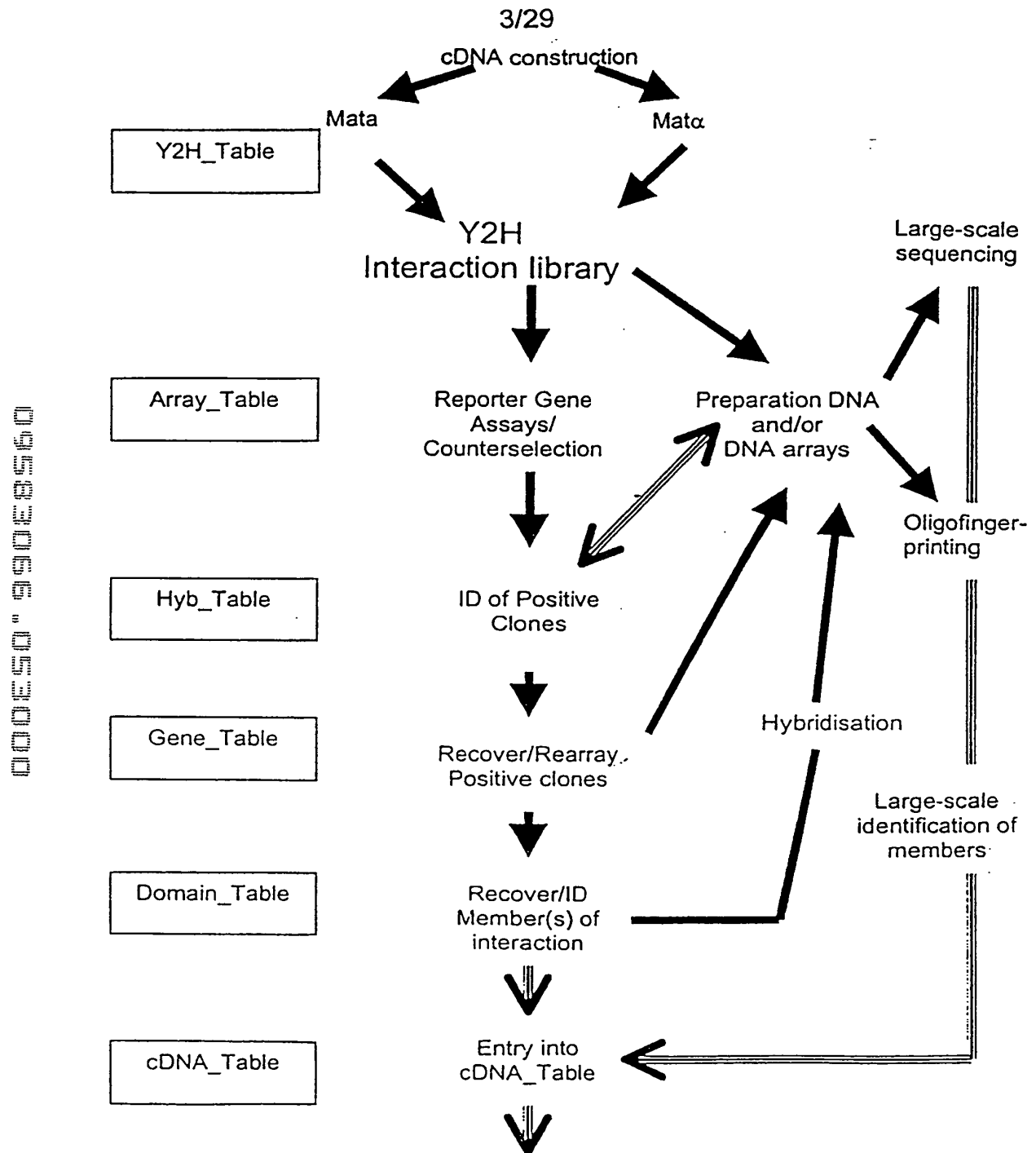


Figure 2



(continuation from the previous page)

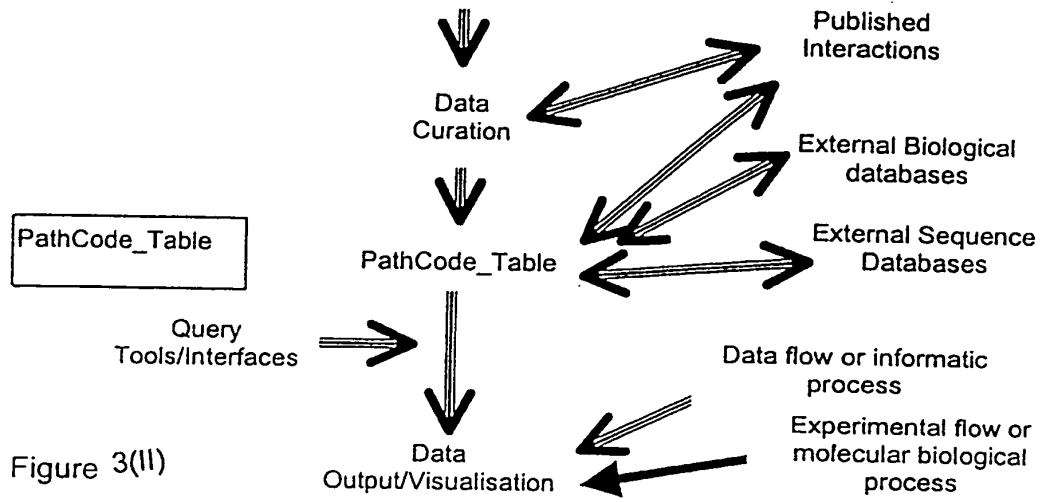


Figure 3(II)

00050" 990E8560

000050" 99000000

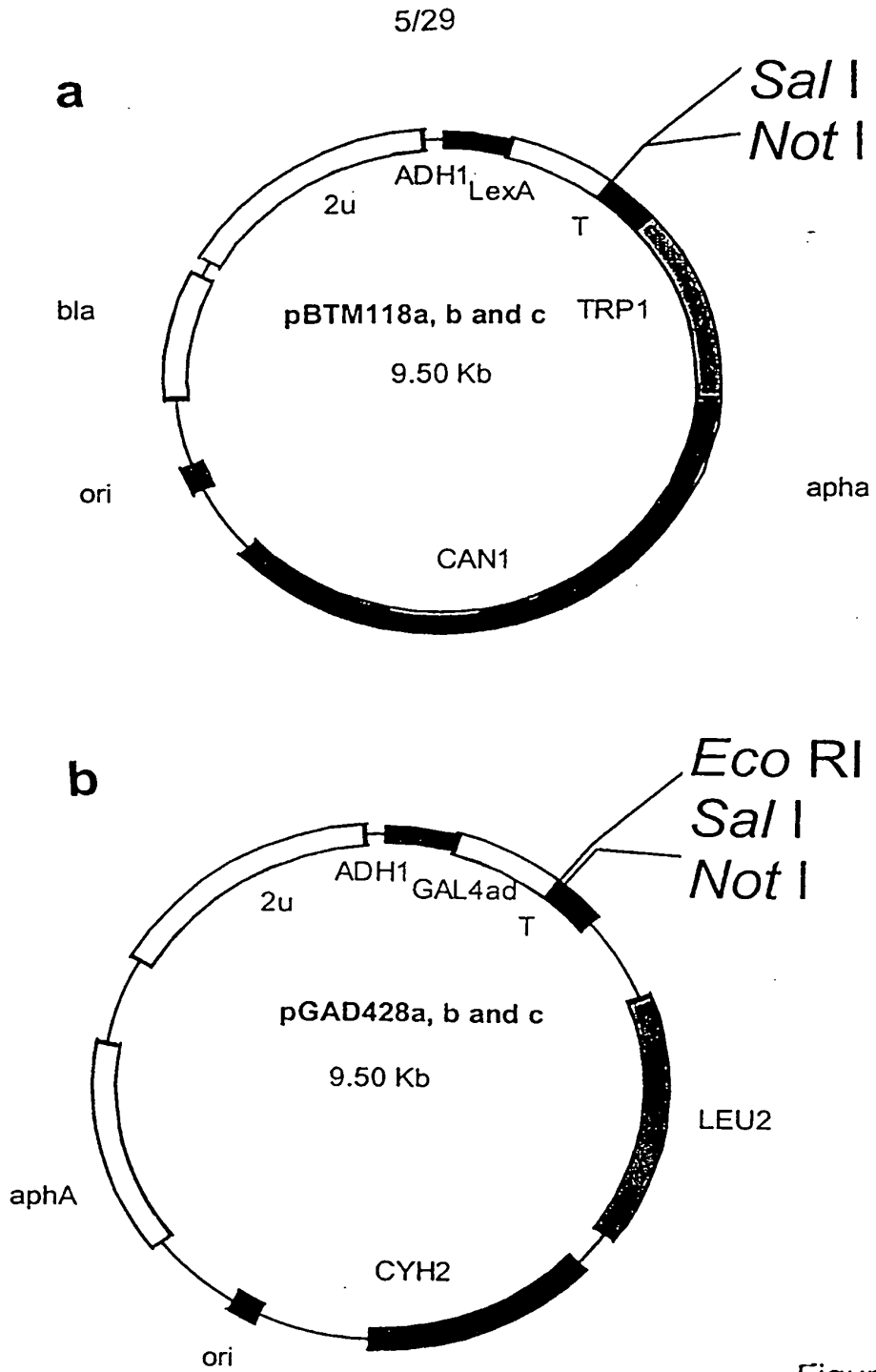


Figure 4 (I)

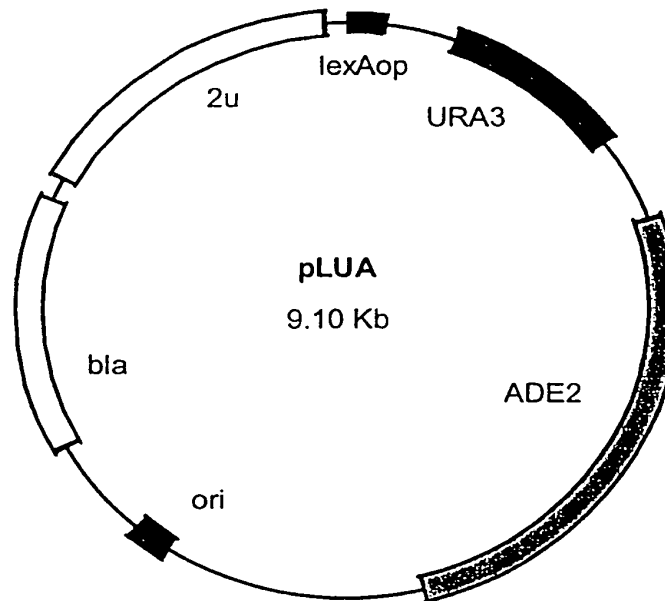
b

a) TCG AGT CGA OGC GGC CGC TAA CCG G  
*Sal* I *Not* I STOP

b) TCG AGG TCG ACG CGG CCG CAG TAA CCG G  
*Sal* I *Not* I STOP

c) TCG AGA GTC GAC GCG GCC GCT TAA CCG G  
*Sal* I *Not* I STOP

Figure 4 (II) ↑  
 Figure 5 ↓



000050" 99000000

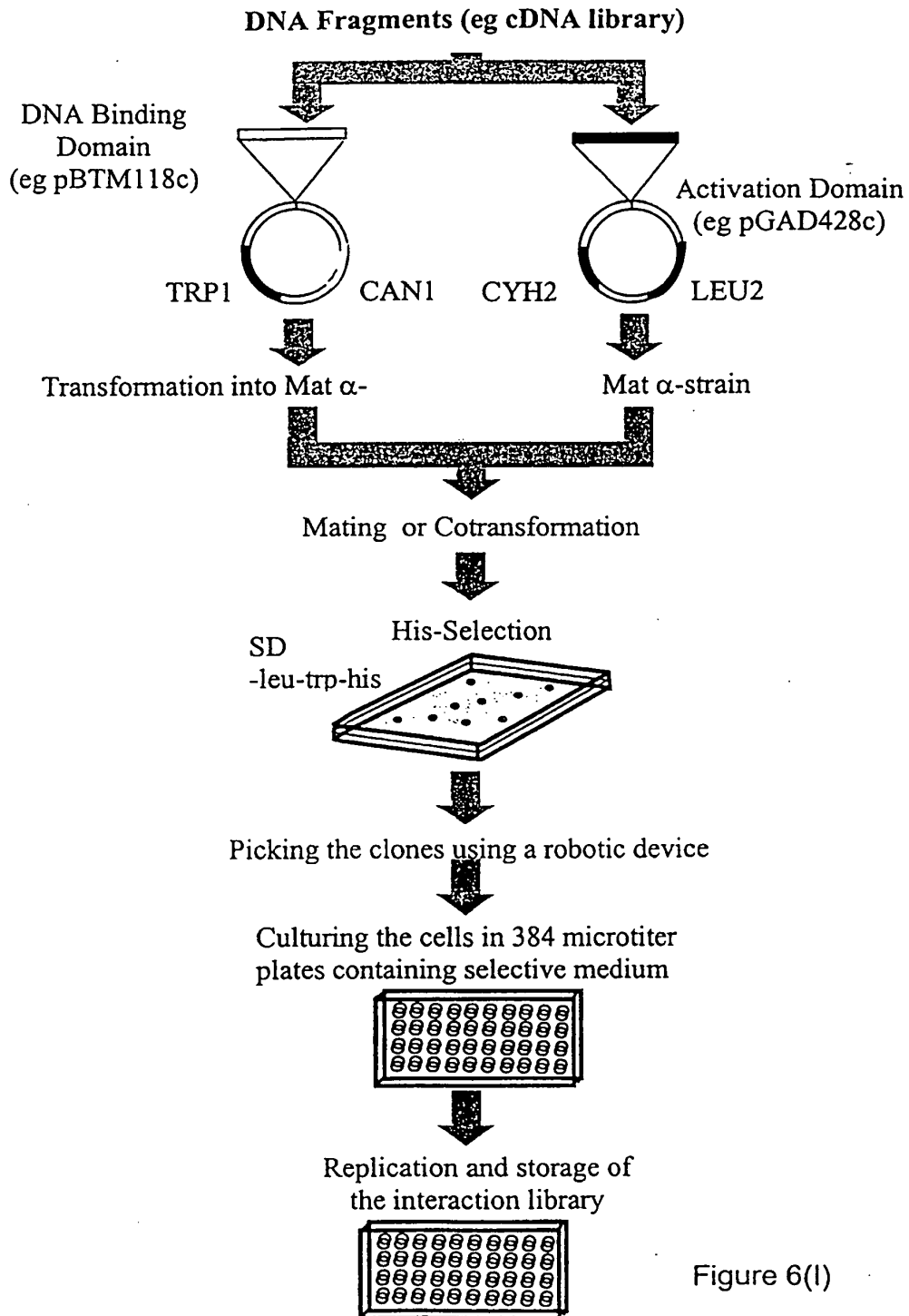
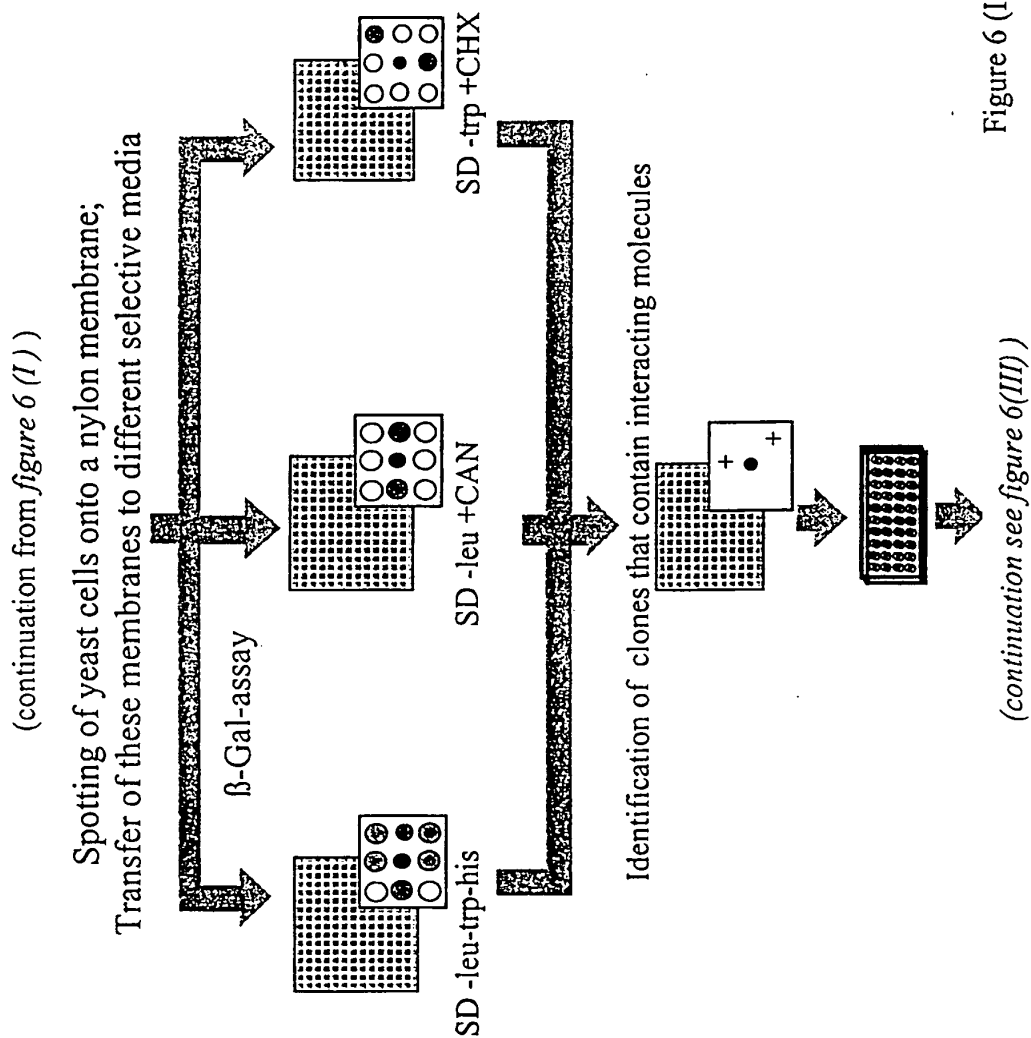


Figure 6(l)

000000" 9900000000

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# Identification of members comprising interactions

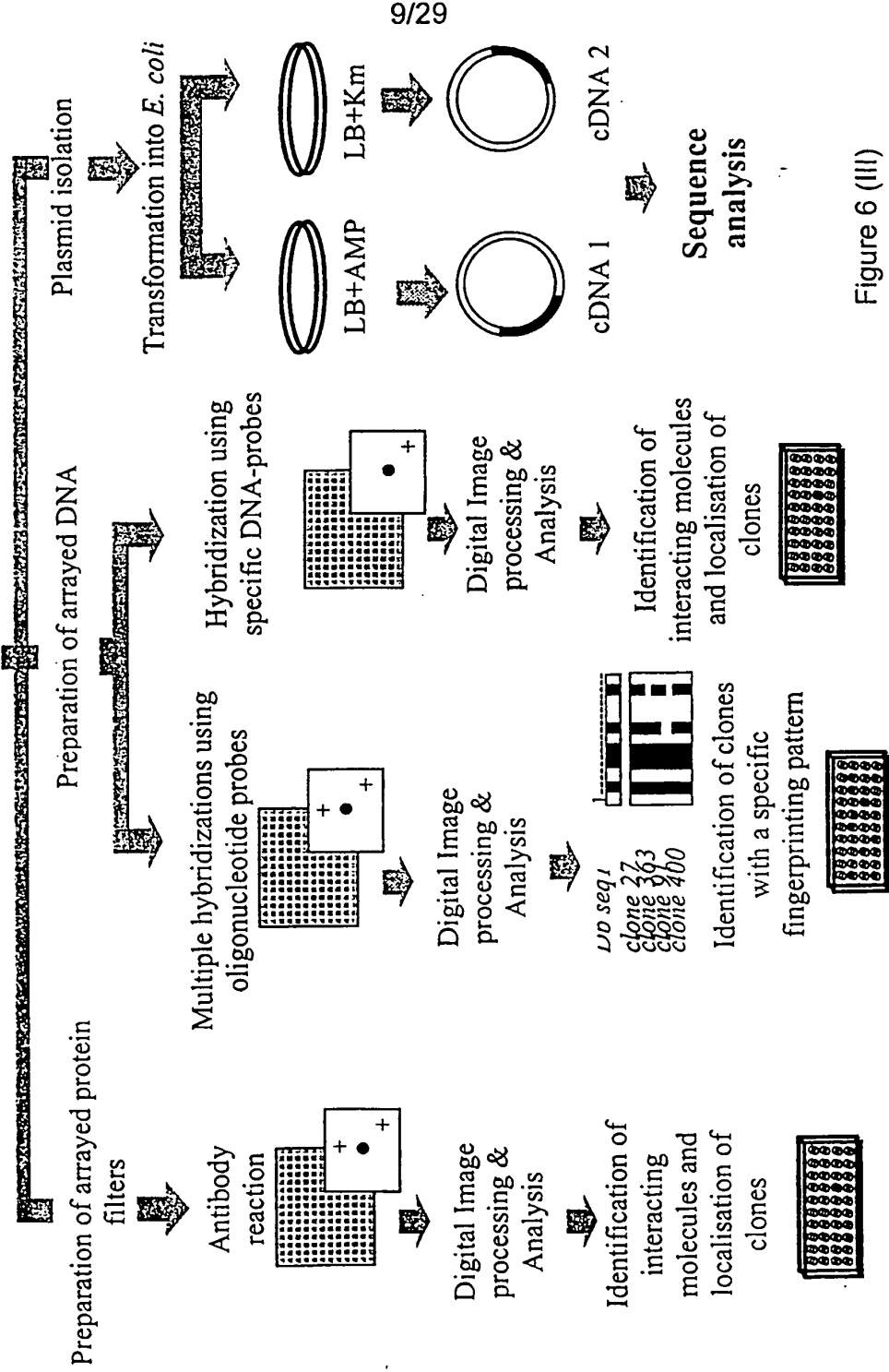
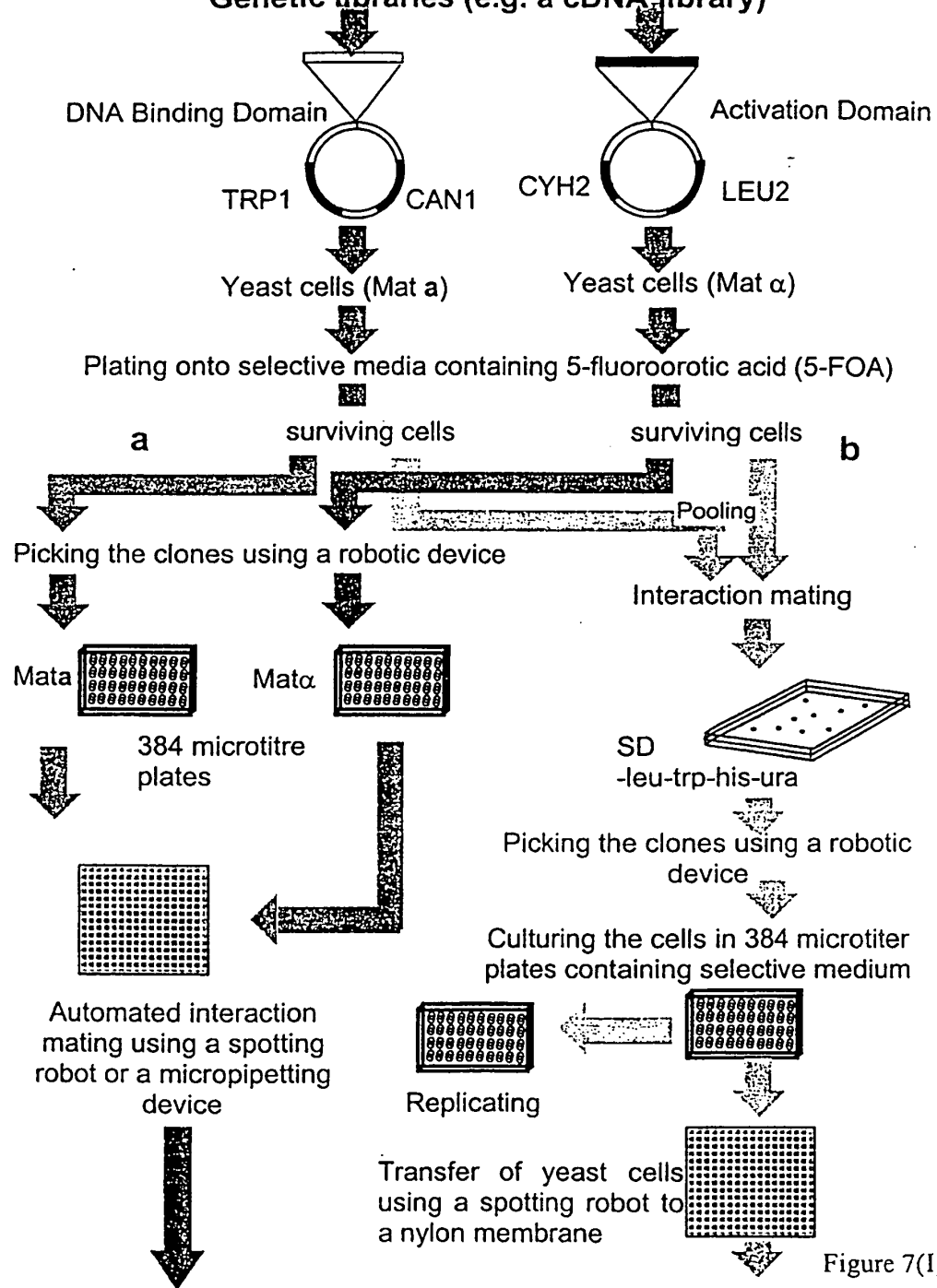


Figure 6 (III)

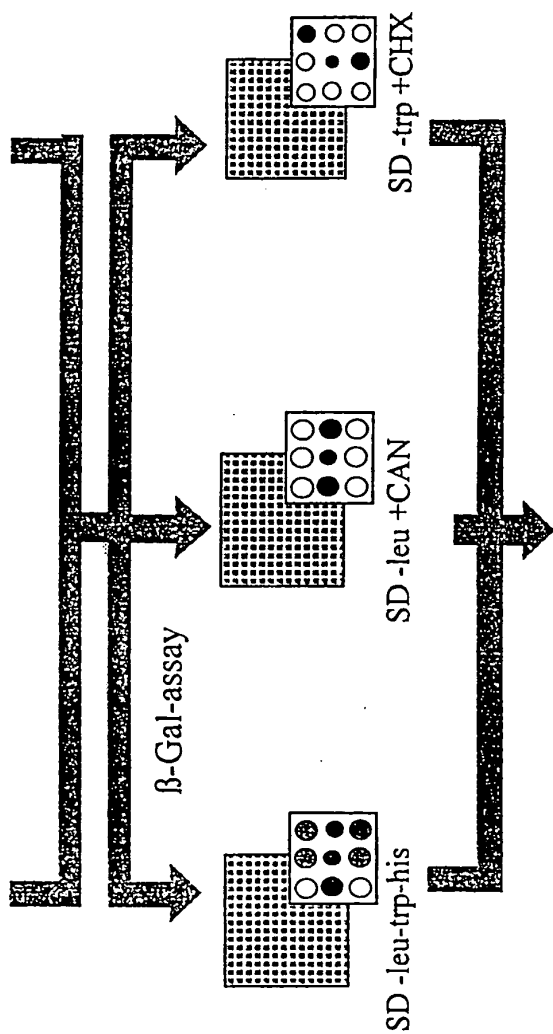
10/29  
Genetic libraries (e.g. a cDNA library)



09583066 053000

000E50" 990E8560

(continuation from figure 7 (I) )



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Figure 7 (II)

(continuation see figure 7(II) )

# Identification of genetic elements comprising interactions

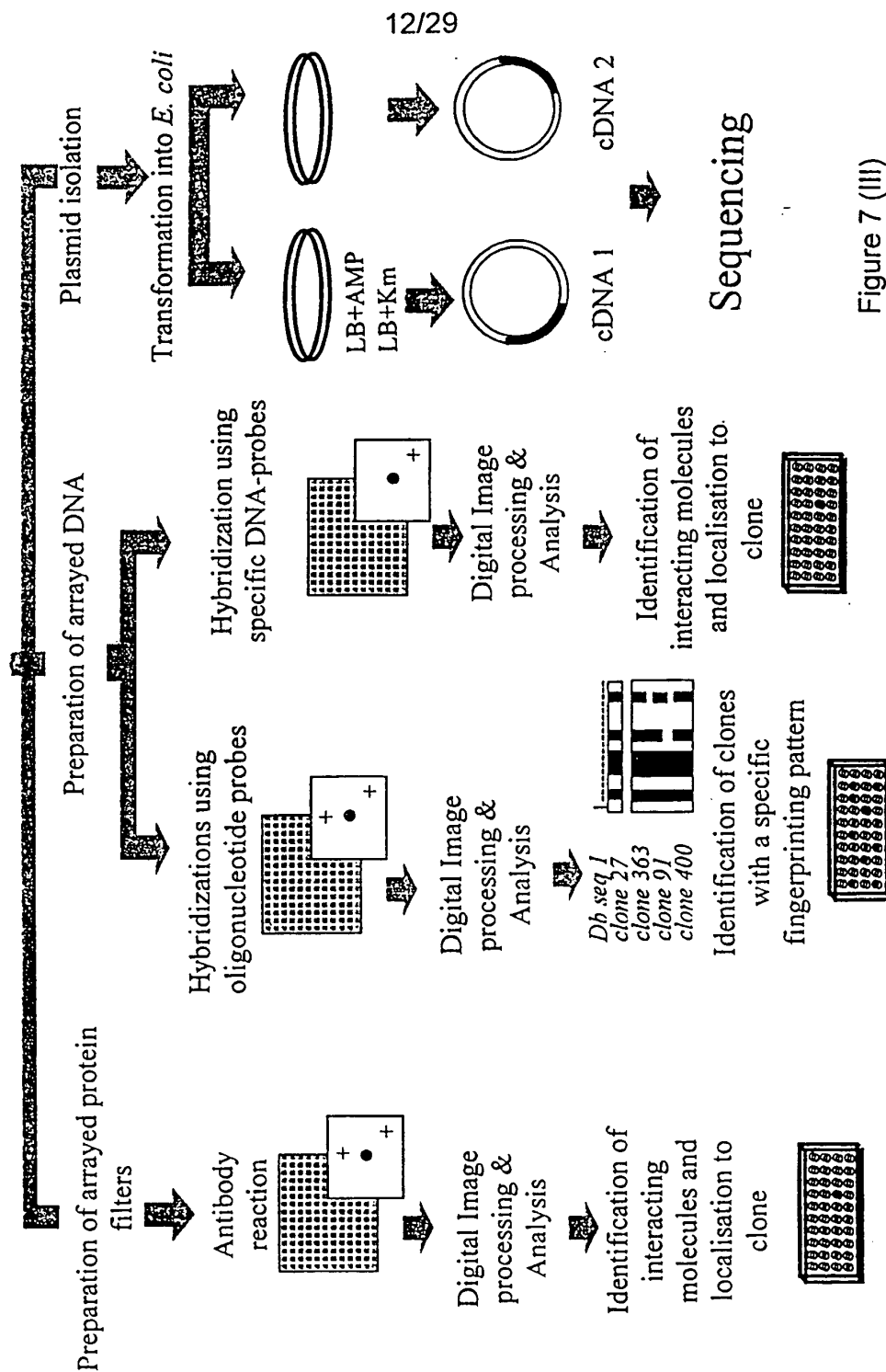


Figure 7 (III)

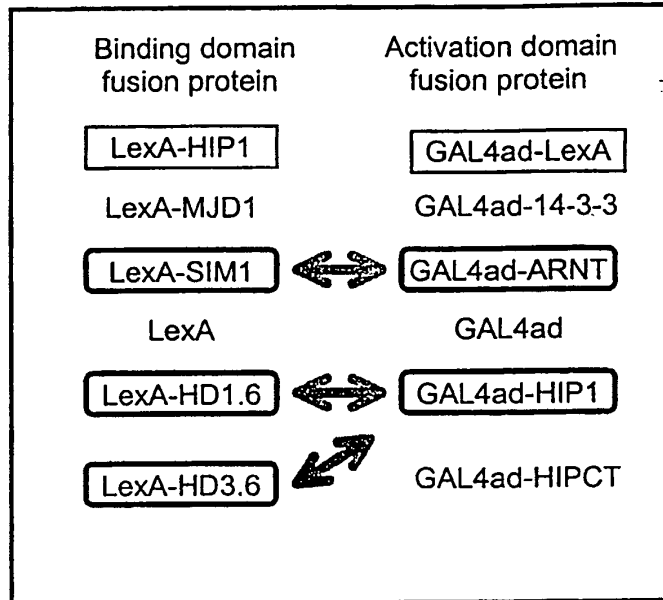


Figure 8

09583066.053000

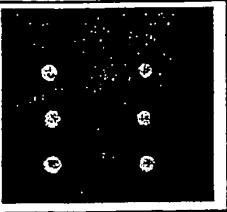
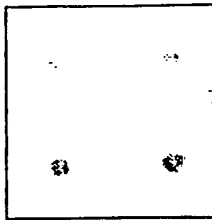
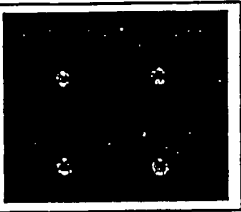
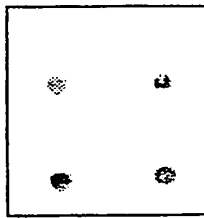

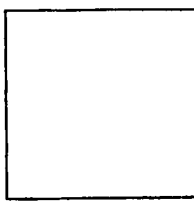

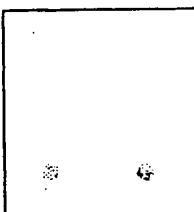
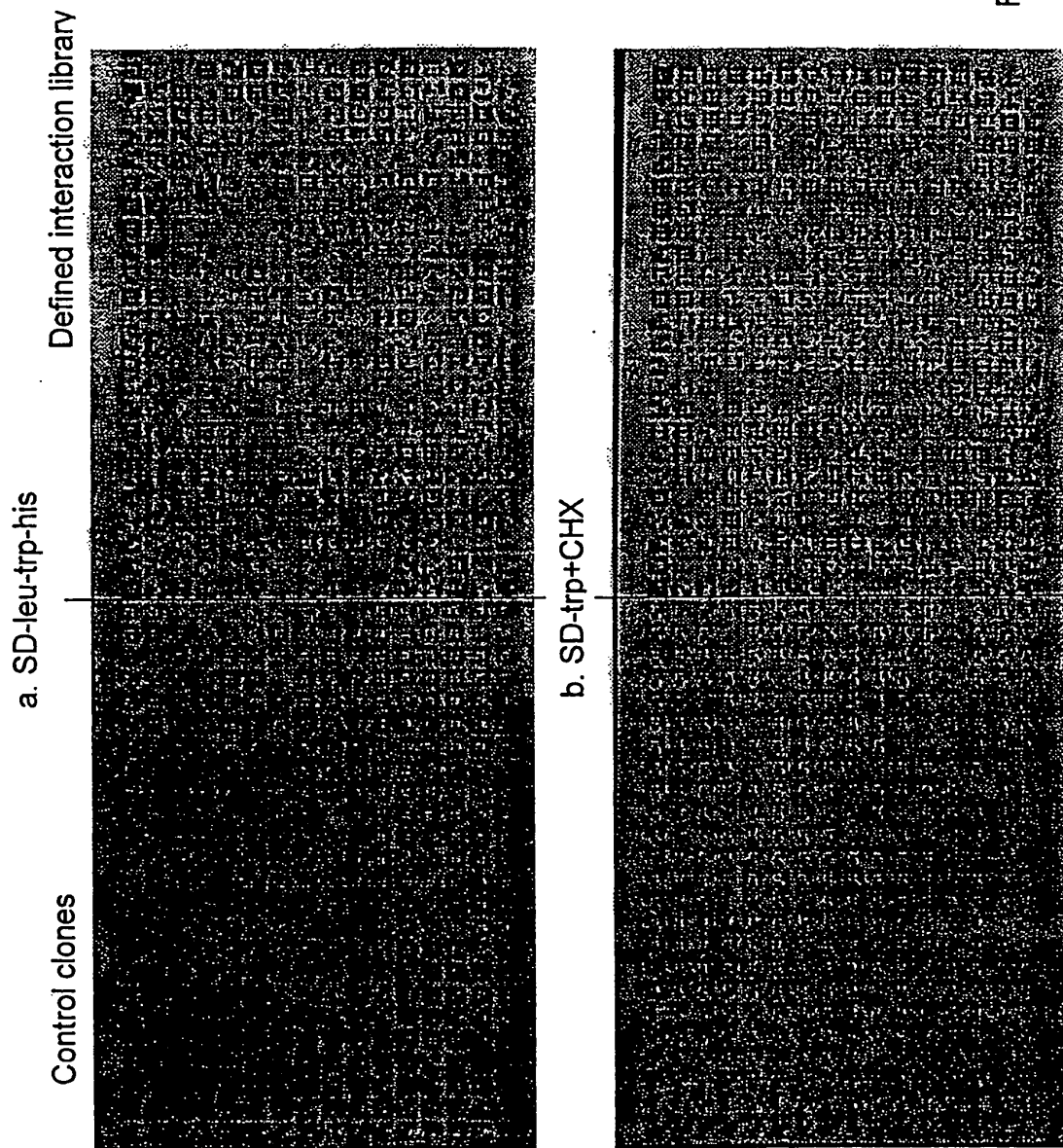
growth	$\beta$ -gal assay
<p>SD-leu-trp</p> 	
<p>SD-leu-trp-his</p> 	
<p>SD-leu+CAN</p> 	
<p>SD-trp+CHX</p> 	

Figure 9

000E50" 990E8560



000E50" 990E8560

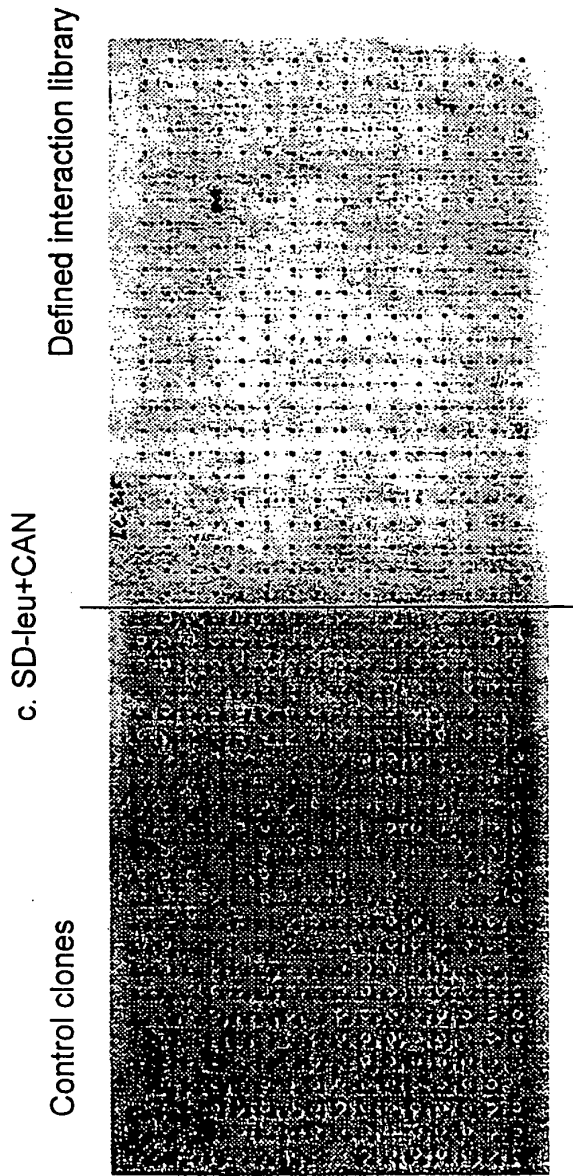


Figure 10 (II)



000E50" 990E8560

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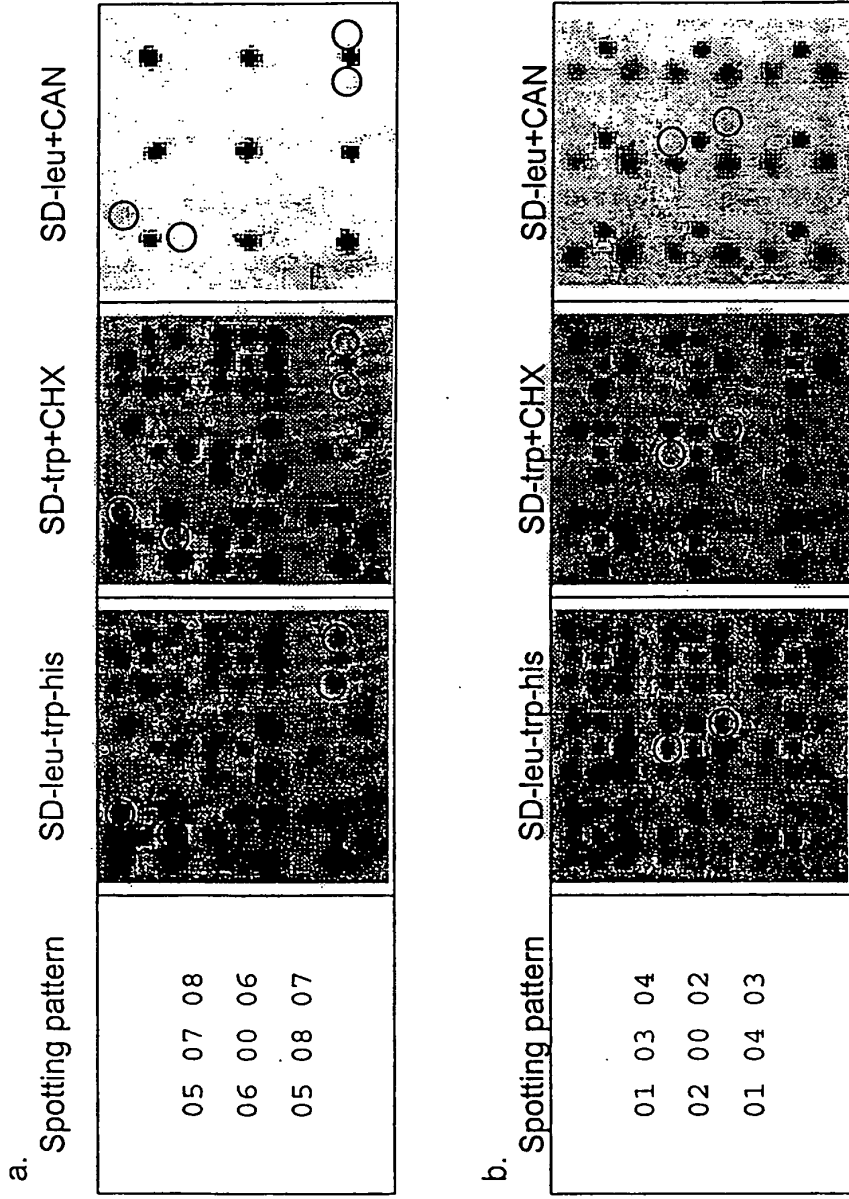


Figure 11

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GPC - PositiveFind						<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>
	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN				
675	208	2	0	0				
687	2022	2	0	0				
689	2P1	3	0	0				
690	2P2	2	0	0				
693	2P5	3	0	0				
696	2P8	3	0	0				
699	2P12	3	0	0				
709	2P23	2	0	0				
710	2P24	2	0	0				
711	3A2	3	0	0				
712	3A3	3	0	0				
713	3A4	3	0	0				
714	3A5	3	0	0				
716	3A7	3	0	0				
717	3A8	3	0	0				
718	3A11	3	0	0				
719	3A12	2	0	0				
720	3A14	3	0	0				
721	3A15	3	0	0				
722	3A18	2	0	0				
723	3A21	2	0	0				
724	3A23	2	0	0				
725	3A24	3	0	0				
727	3B2	3	0	0				
728	3B3	3	0	0				
729	3B4	3	0	0				
730	3B5	3	0	0				
731	3B6	3	0	0				
732	3B7	3	0	0				
733	3B8	3	0	0				
734	3B9	3	0	0				
735	3B10	3	0	0				
736	3B12	3	0	0				
737	3B13	3	0	0				
738	3B14	3	0	0				
739	3B15	3	0	0				
741	3B17	2	0	0				

b.

GPC - PositiveFind				
1	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN
1	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN
1733	6L22	3	0	0
2302	8N24	2	0	0

Figure 12

**0963071**

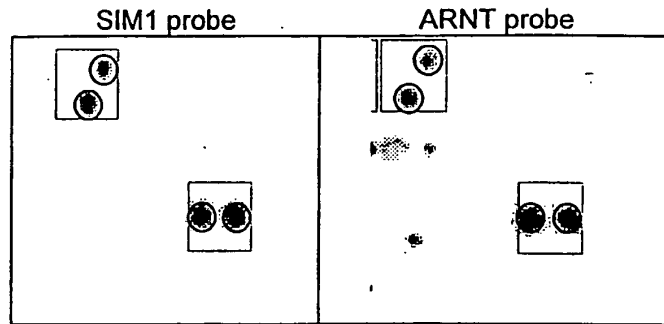


Figure 13

000E50" 990E8560



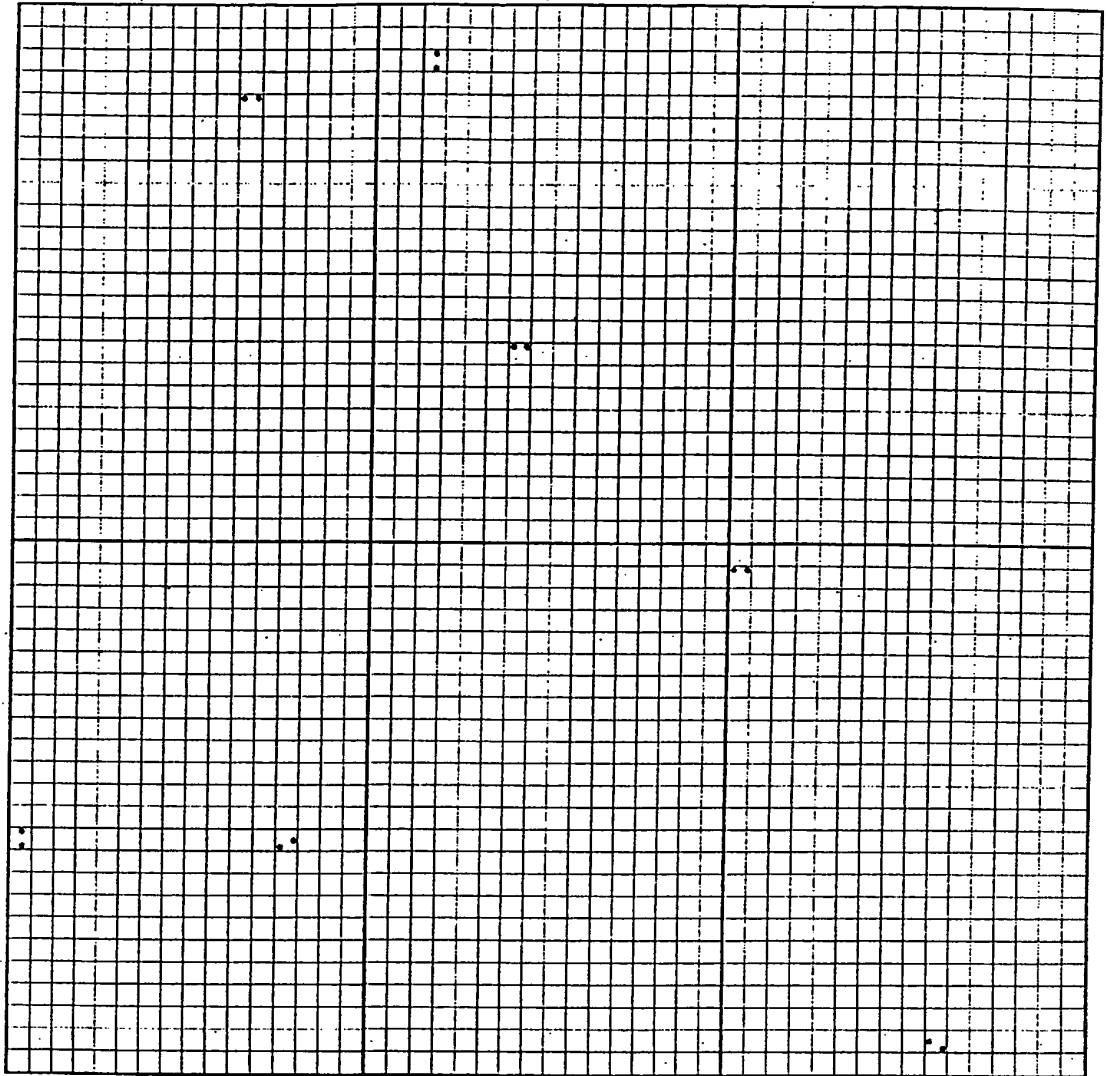
000E50" 990E8560



Figure 15



09583065, 053000



1st Hybridisation

Protein A: ●

Interaction Map

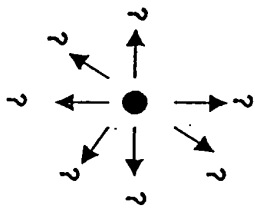
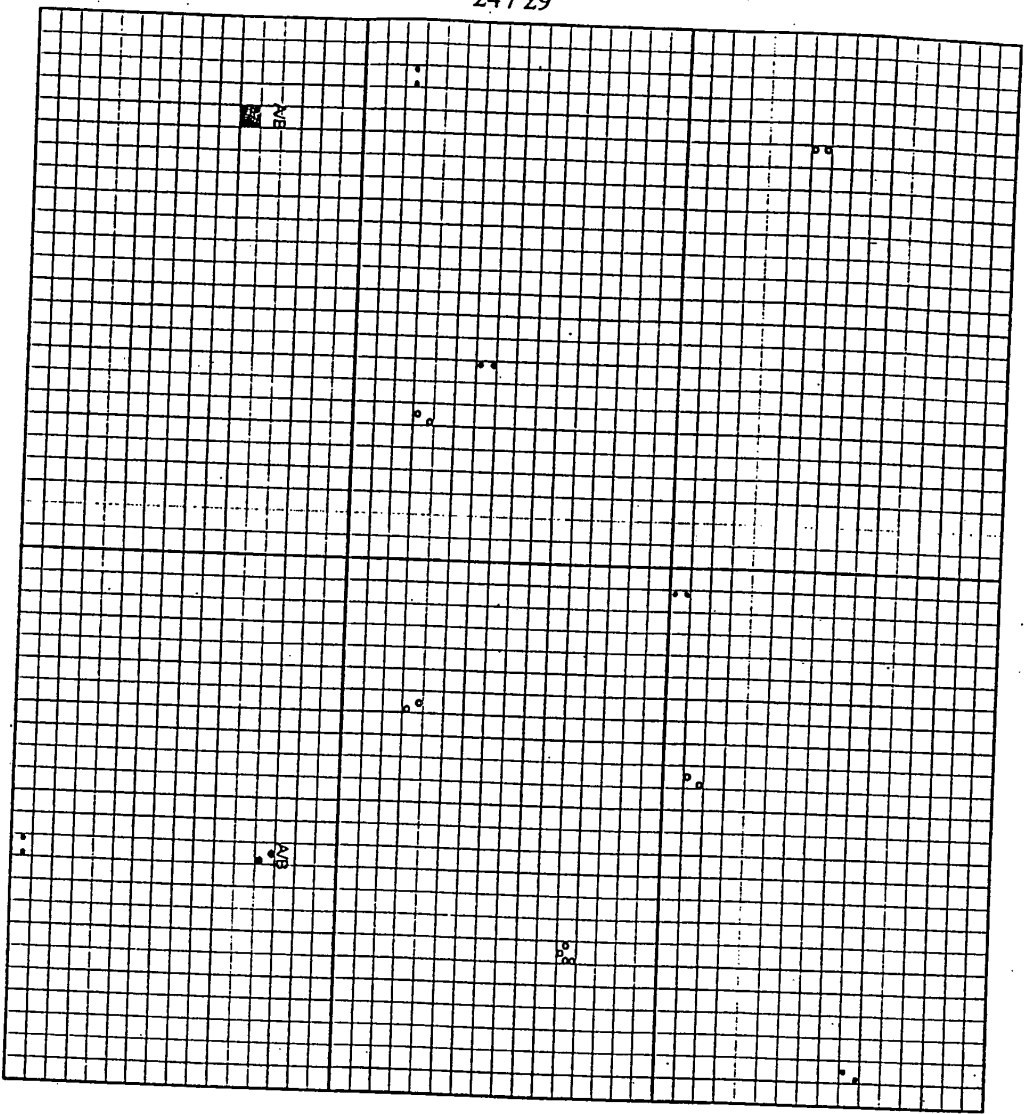


Figure 17



1st Hybridisation

Protein A: ●

2nd Hybridisation

Protein B: ○

Interaction Map

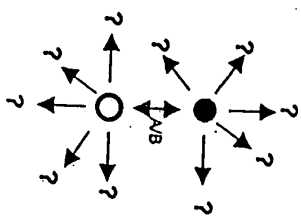


Figure 18

09583055 053000



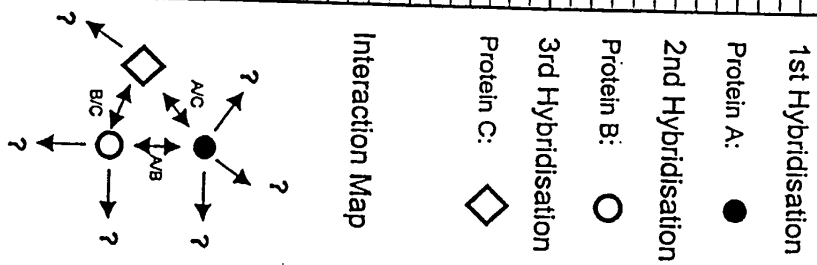
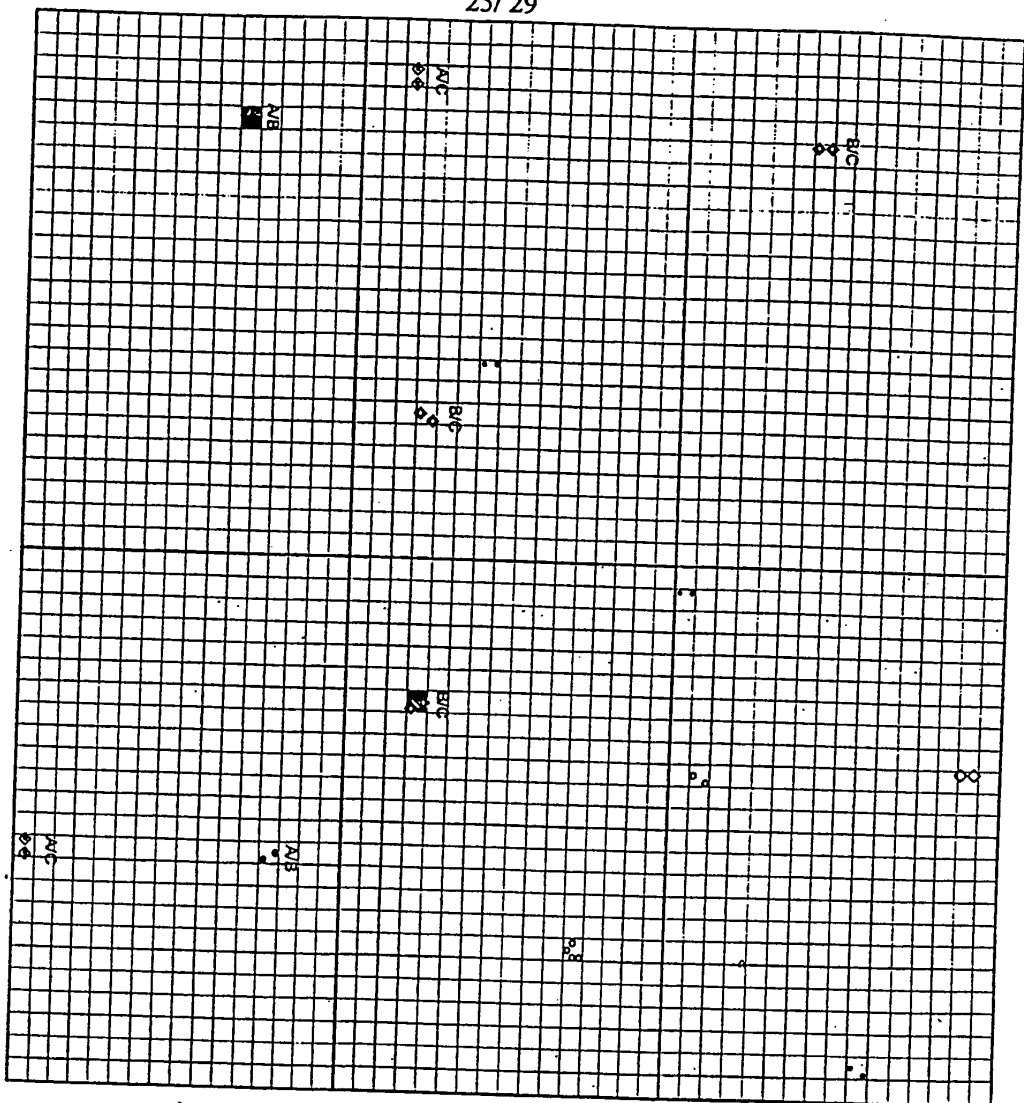


Figure 19

09583066.053000

000E50" 990E8560

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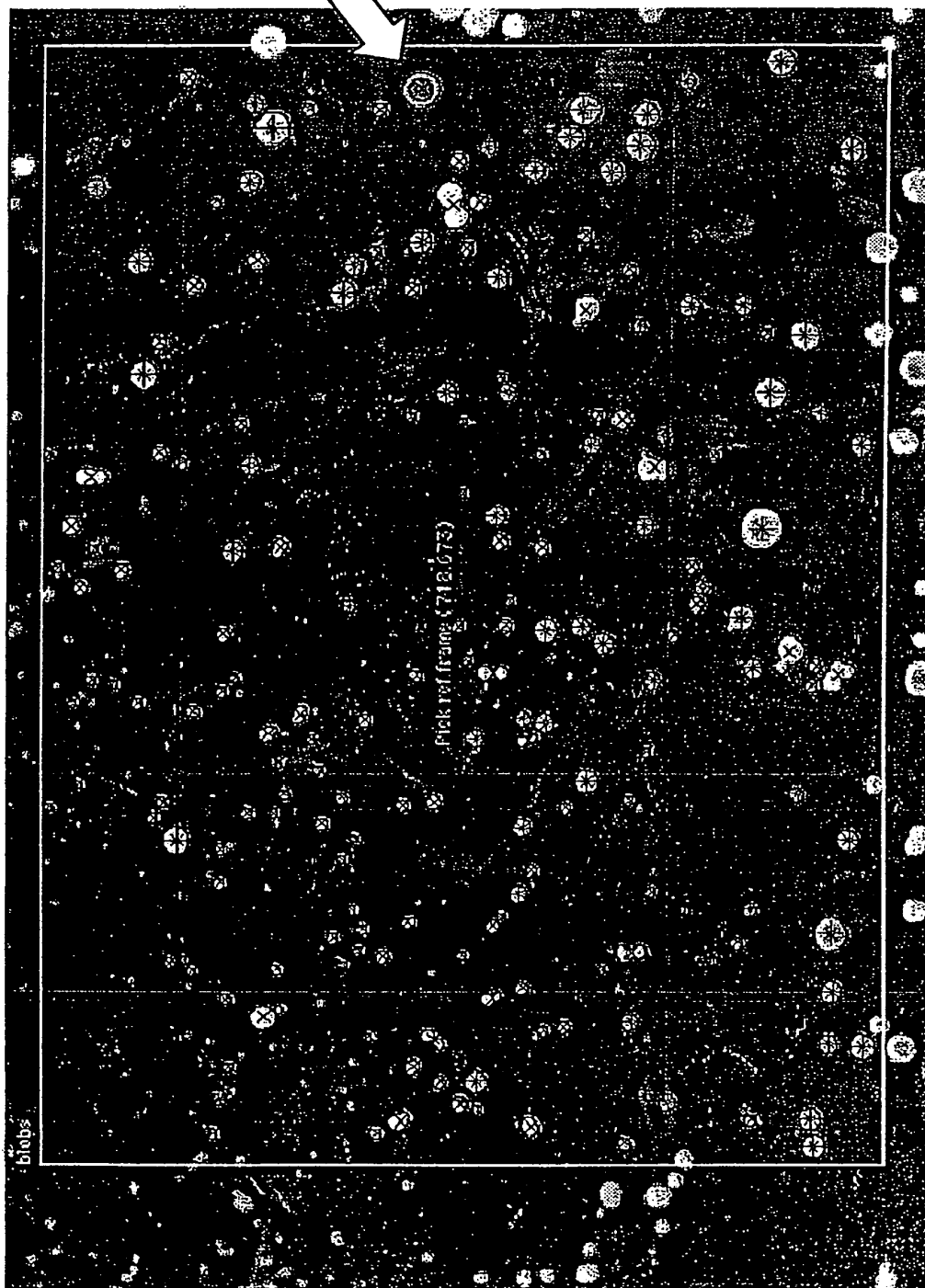


Figure 20

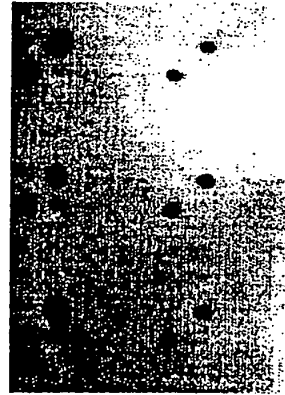
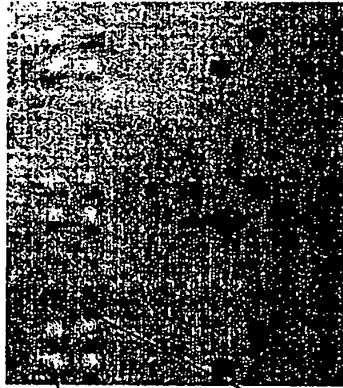
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YPD

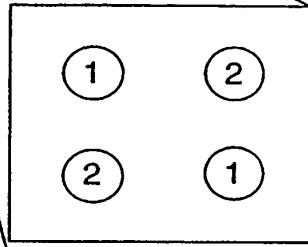
X-Gal

b.

c.



a.



2 mm

000050" 990E8560

Figure 21

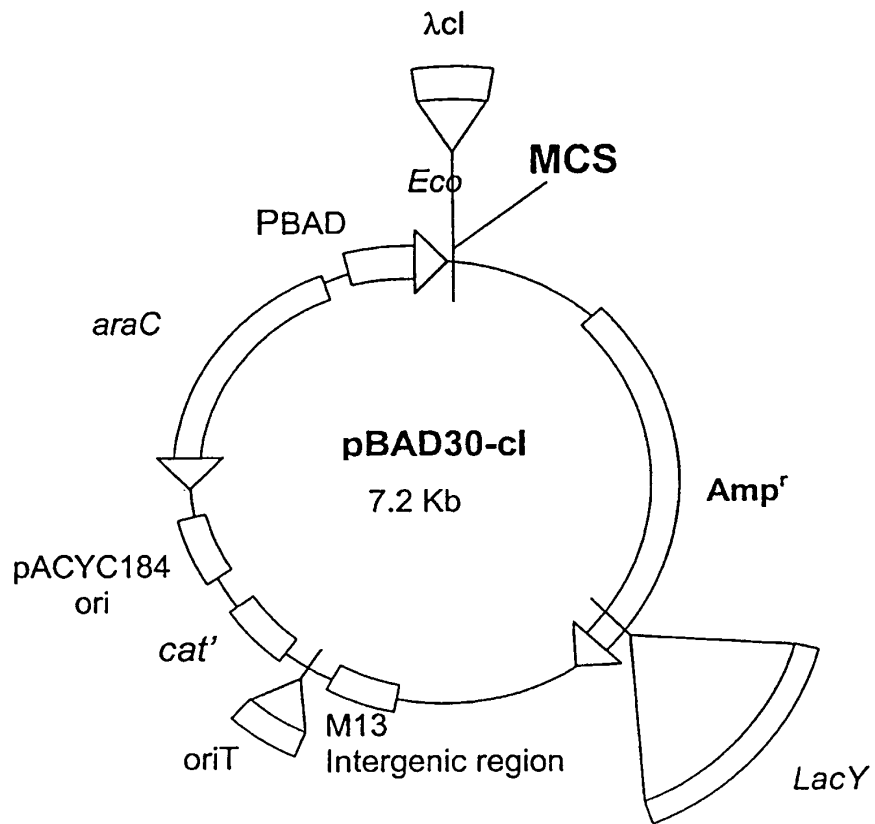


Figure 22 (I)

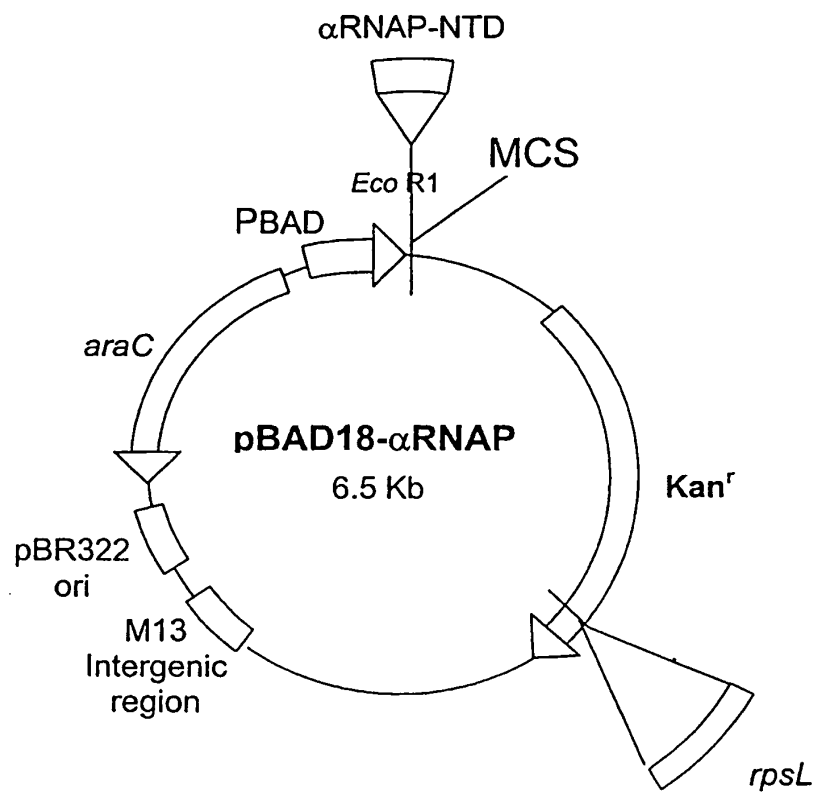


Figure 22 (II)

000550" 990E8560